SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANTS: Robert P. Kimberly

(ii) TITLE OF INVENTION: GENETIC POLYMORPHISM IN THE RECEPTOR FOR IgA

(iii) NUMBER OF SEQUENCES: 4

(iv) CORRESPONDENCE ADDRESS:

(A)	ADDRESSEE:	Ellen S. Cogen Gifford, Krass, Groh, Sprinkle, Anderson & Citkowski, P.C.
(B)	STREET:	280 N. Old Woodward Avenue Suite 400
(C)	CITY:	Birmingham
(D)	STATE:	Michigan
(E)	COUNTRY:	U.S.
(F)	ZIP:	48009

(v) COMPUTER READABLE FORM:

(A)	MEDIUM TYPE:	Floppy disk
(B)	COMPUTER:	IBM
(C)	OPERATING SYSTEM:	Windows 98
(D)	SOFTWARE:	WordPerfect 7.0

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Ellen S. Cogen
(B) REGISTRATION NUMBER: 38,109
(C) REFERENCE/DOCKET NUMBER: UAB-14252/22

(ix) TELECOMMUNICATION INFORMATION:

(A)	TELEPHONE:	248-647-6000
(B)	TELEFAX:	248-647-5210

(2) INFORMATION FOR SEQ ID NO. 1:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 39
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE:
(A) Description: DNA (genomic)

(iii) HYPOTHETICAL:

(iv) ANTISENSE:

(vi) ORIGINAL SOURCE:
(B) STRAIN:
(C) INDIVIDUAL ISOLATE:
(D) DEVELOPMENTAL STAGE:
(F) TISSUE TYPE:
(G) CELL TYPE:
(H) CELL LINE:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO. 1:

TGT AAA ACG ACG GCC AGT AGC ACG ATG GAC CCC AAA CAG

(3) INFORMATION FOR SEQ ID NO. 2:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 38
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE:
(A) Description: DNA (genomic)

(iii) HYPOTHETICAL:

(iv) ANTISENSE:

(vi) ORIGINAL SOURCE:
(B) STRAIN:

(C) INDIVIDUAL ISOLATE:
(D) DEVELOPMENTAL STAGE:
(F) TISSUE TYPE:
(G) CELL TYPE:
(H) CELL LINE:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO. 2:

CAG GAA ACA GCT ATG ACC GGT GTT CCC CAC TTT GGT GC

(4) INFORMATION FOR SEQ ID NO. 3:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 38
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE:
(A) Description: DNA (genomic)

(iii) HYPOTHETICAL:

(iv) ANTISENSE:

(vi) ORIGINAL SOURCE:
(B) STRAIN:
(C) INDIVIDUAL ISOLATE:
(D) DEVELOPMENTAL STAGE:
(F) TISSUE TYPE:
(G) CELL TYPE:
(H) CELL LINE:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO. 3:

TGT AAA ACG ACG GCC AGT AGA ATA TTT CCC TCA CGT GC

(5) INFORMATION FOR SEQ ID NO. 4:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 39
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE:
(A) Description: DNA (genomic)

(iii) HYPOTHETICAL:

(iv) ANTISENSE:

(vi) ORIGINAL SOURCE:
(B) STRAIN:
(C) INDIVIDUAL ISOLATE:
(D) DEVELOPMENTAL STAGE:
(F) TISSUE TYPE:
(G) CELL TYPE:
(H) CELL LINE:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO. 4:

CAG GAA ACA GCT ATG ACC CTG GCT CCT CTC TGC CTT CAC